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Genome editing of avian species: implications for animal use and welfare

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Abstract

Avian species are used as model systems in research and have contributed to ground-breaking concepts in developmental biology, immunology, genetics, virology, cancer and cell biology. The chicken in particular is an important research model and an agricultural animal as a major contributor to animal protein resources for the global population. The development of genome editing methods, including CRISPR/Cas9, to mediate germline engineering of the avian genome will have important applications in biomedical, agricultural and biotechnological activities. Notably, these precise genome editing tools have the potential to enhance avian health and productivity by identifying and validating beneficial genetic variants in bird populations. Here, we present a concise description of the existing methods and current applications of the genome editing tools in bird species, focused on chickens, with attention on animal use and welfare issues for each of the techniques presented.

Keywords

Poultry, chicken, primordial germ cells, poultry welfare, genome editing, CRISPR/Cas9

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Introduction

Genome editing (GE) is a method for the rapid introduction of precise changes into an organism's genome. GE tools consist of programmable site-specific nucleases, that is meganucleases, zinc finger nucleases, TALENs, and CRISPR/Cas9.^{1,2} All of these programmable site-specific nucleases or 'genome editors' can be used to efficiently create precise genetic changes, generated through double-stranded breaks (DSBs) at specific locations in the genome. The DSBs are repaired by two conserved cellular machinery pathways: the non-homologous end-joining (NHEJ) pathway or the rarer homology-directed repair (HDR) pathway. The NHEJ pathway is often used by the cell to repair DNA damage and can result in small genetic insertions or deletions (INDELs) at the DSB. The more accurate HDR pathway occurs in the presence of a sister chromatid or an exogenous DNA fragment containing a homologous region spanning the DSB which are used as templates to repair the DSB.³ The HDR repair process has been exploited to generate defined small genetic deletions, insertions, single base-pair changes in a gene, and even the directed integration of large

exogenous transgenes precisely into the host cellular genome. What is novel is that this technology enables the creation of site-specific genetic changes without leaving any other modifications in the genome (footprintless editing), and these changes are indistinguishable from naturally occurring genetic variants. The development of genome targeting through HDR has significantly improved 'transgenic' research, by creating both plant and animal models with more precise and defined manipulations of the genome.

Avians serve as an important source of animal protein, and as a research model for the study of developmental biology, immunology and infectious diseases.⁴ The chicken is both the principal avian research model and an agricultural animal with a global population

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numbering over 60 billion animals annually. Commercial chicken flocks are the source of fertile eggs for many experiments and these eggs can also be for public food consumption. Specialised chicken research flocks are also maintained at research institutes in compliance with the Animals (Scientific Procedures) Act 1986 in the UK and these flocks have been selectively bred to have defined genetic resistances and susceptibilities to avian diseases, specific developmental genetic mutations, or have been genetically modified to carry exogenous transgenes. Many other bird species such as quail, duck and zebra finch are also used as standard laboratory research models.⁵ The annual Statistics of Scientific Procedures on Living Animals Great Britain 2019 reported 0.13 million avians (8%) among all animal groups were used in experimental procedures, whilst only 0.1% of birds were used for the creation and breeding of genetically altered (GA) animals.⁶

Avian flocks are susceptible to a plethora of bacterial and viral pathogens threatening their health and welfare, and directly threatening flock security and the global agro-economy.⁷ Bacterial pathogens such as *Campylobacter jejuni* and *Salmonella enteritidis* cause health risks from raw poultry products and consequently cause foodborne illness in humans.⁸ Viral pathogens causing Newcastle Disease and Marek's Disease are impactful on both wild birds and the poultry industry⁹ and the avian influenza A virus (IAV) creates both animal and public health concerns.¹⁰ GE avian models to investigate disease-resistance are of special interest to the poultry industry. GE avian research models to address key unanswered questions in developmental biology, immunology, physiology, cell biology and neural biology are also important to life scientists. Chicken eggs have also been used for the production of biopharmaceutical products.¹² This review will focus on advances of GE tools in the chicken, which is the most developed avian research species, and highlight advances made in other avian species.

GE of bird species

Mammalian assisted *in vitro* reproductive technologies are well developed and mimic *in vivo* reproductive physiology. They are based on synchronisation and super-ovulation of the donor animal followed by oocyte retrieval, *in vitro* fertilisation or cloning and subsequent re-implantation into a surrogate host female (Figure 1 (a)). The one-cell fertilised zygote serves as an accessible recipient for the microinjection of genome editors to create specific genetic alterations. Gordon and Ruddle first reported the pronuclear microinjection method by directly delivered the exogenous DNA into the nucleus of the zygote.¹³ Pronuclear

microinjection leads to the random integration of the exogenous DNA that can result in high mosaicism and aberrant transgene expression. Scientists spent decades refining and adopting this technology to other mammalian species such as rabbit, pig, sheep, goat and cattle.^{14–16} Genome editors can similarly be introduced into the cytoplasm or the pronuclei of the zygote. The genetic changes created in the early stage embryo are carried to the successive developmental stages to generate mosaic founder animal that will contain the genetic change in some or all of its cells, including the reproductive germ cells. Targeted genome edited somatic cells can also be used as a nuclear donor in enucleated oocytes in the process of somatic cell nuclear transfer (SCNT) or hand-made cloning^{17,18} to produce live offspring carrying the desired genome edit in all cells of the animal. This two-step protocol, editing of somatic cells followed by SCNT, has proven to be efficient and highly precise to create specific alterations to the livestock genome in a single generation without the need for breeding multiple generations to obtain homozygous animals containing the desired edit. However, limitations of this technique include the difficulty in targeting somatic cells, low cloning efficiencies and large offspring syndrome with its associated birth complications.¹⁹ Even with these caveats, genome editing using SCNT has been accomplished in pig, cattle, goat and sheep species.^{20–23}

In contrast to mammalian species, avians have a unique reproductive physiology as well as distinctive structure of the ovum and pre-gastrulation stage embryo. In birds, the macrolecithal ovum is released from the ovary surrounded by a tough vitelline membrane, which serves as a protective layer around the deposited yolk. The ovulated ovum is rapidly fertilised with sperm that is stored by the female in sperm glands located in the oviduct. The single-cell zygote consists of a small pool of cytoplasm containing the fused pronuclei on the surface of the large yolk mass. The zygote takes approximately 24 h to pass through the oviduct region; the egg white is first added to enclose the yolk, followed by a shell membrane and a hard shell that are added during the final stages of passage through the oviduct. During this journey of the zygote from infundibulum to cloaca for laying of the egg, the single-cell zygote has undergone multiple rounds of cleavage and the pool of cytoplasm has developed into the blastoderm layer consisting of many tens of thousands of undifferentiated cells. Subsequently, the laid egg can be artificially or naturally incubated until hatching (Figure 1(b)). Thus, the relatively simple straightforward one-cell embryo microinjection procedures of generating genome edited mammalian embryos or SCNT cannot be easily replicated in birds as the early stage embryo is not accessible. Recent advances,

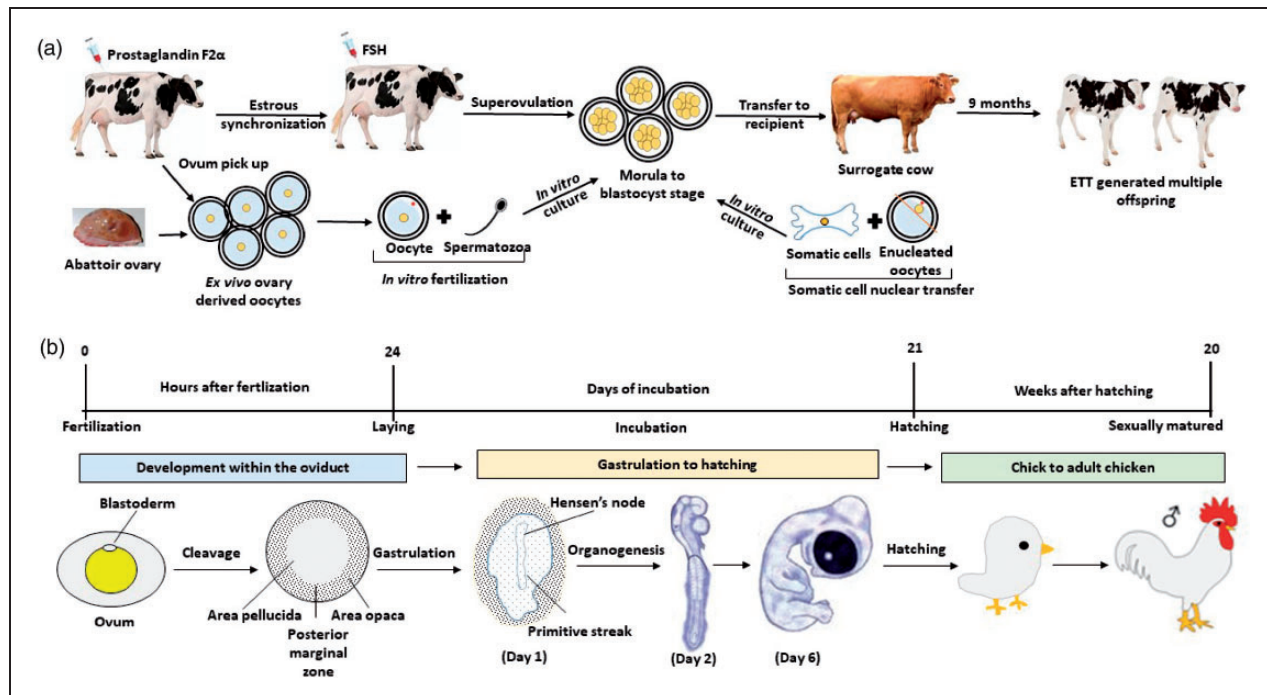


Figure 1. Schematic diagram of the reproductive physiology of bovine and chicken species. (a) Production of high genetic merit calves through the process of the synchronisation, superovulation, ovum pick up, *in vitro* fertilization or cloning, and embryo transfer to the surrogate cow. (b) Reproductive cycle of chicken through the process of natural mating or artificial insemination followed by incubation, hatching and raising to sexual maturity.

however, have been made manipulating the early stage of ovum using intracytoplasmic sperm injection (ICSI) for fertilisation²⁴ and a surrogate shell for incubation.²⁵ However, this ICSI method has yet to be used for GE as few hatchlings were obtained from the ICSI embryos.

Mouse and human embryonic stem cells (ES cells), derived from the blastocyst stage embryo, are useful cell lines as they can be genetically modified *in vitro* and can be differentiated into any cell type of the body. Mouse ES cells will also contribute to embryonic chimeras and form functional sperm or eggs. Avian ES cells cultured from the avian blastoderm of laid eggs (Figure 1(b)) have been propagated for chicken.²⁶ However, similar to other livestock species, no evidence of germline transmission using cultured (longer than one week) embryonic stem cells has been reported for chicken or other bird species. Instead, chicken ES cells injected into early chicken embryos contributed to somatic tissue but not to the germline after only a short period of *in vitro* culture.^{26,27} This may be because the germ cell lineage, cells that are destined to form sperm and oocytes, are present and segregated as a ~ 50 cell population in the blastoderm of the laid avian egg. This is very different from mammalian species for which the germ cell lineage forms much later in embryonic development. This also signifies that the

delivery of GE tools needs to target this small population of germ cells in avian species in order to achieve genetic transmission to the offspring of the injected animal.

Methods used for CRISPR/Cas9-mediated *in-ovo* GE in birds

In spite of the complex architecture of the avian zygote, researchers have developed several delivery methods to introduce genetic vectors into the early avian embryo and cells from the embryo. As CRISPR/Cas9 genome editors are proving highly efficient at modifying the target species genome, these vectors are revolutionising efforts to manipulate the genome of birds. Current delivery methods to avian embryos and gametes are shown in Figure 2.

Direct electroporation of embryos

The advantage of the avian model is that the developing avian embryo is directly accessible through a hole made in the surrounding egg shell, in a process called 'windowing the egg'. The shell can be resealed and the egg re-incubated until the desired embryonic stage is reached or the embryo can be hatched. Direct *in ovo* electroporations of CRISPR/Cas9 plasmids into the embryo can lead to GE of the electroporated

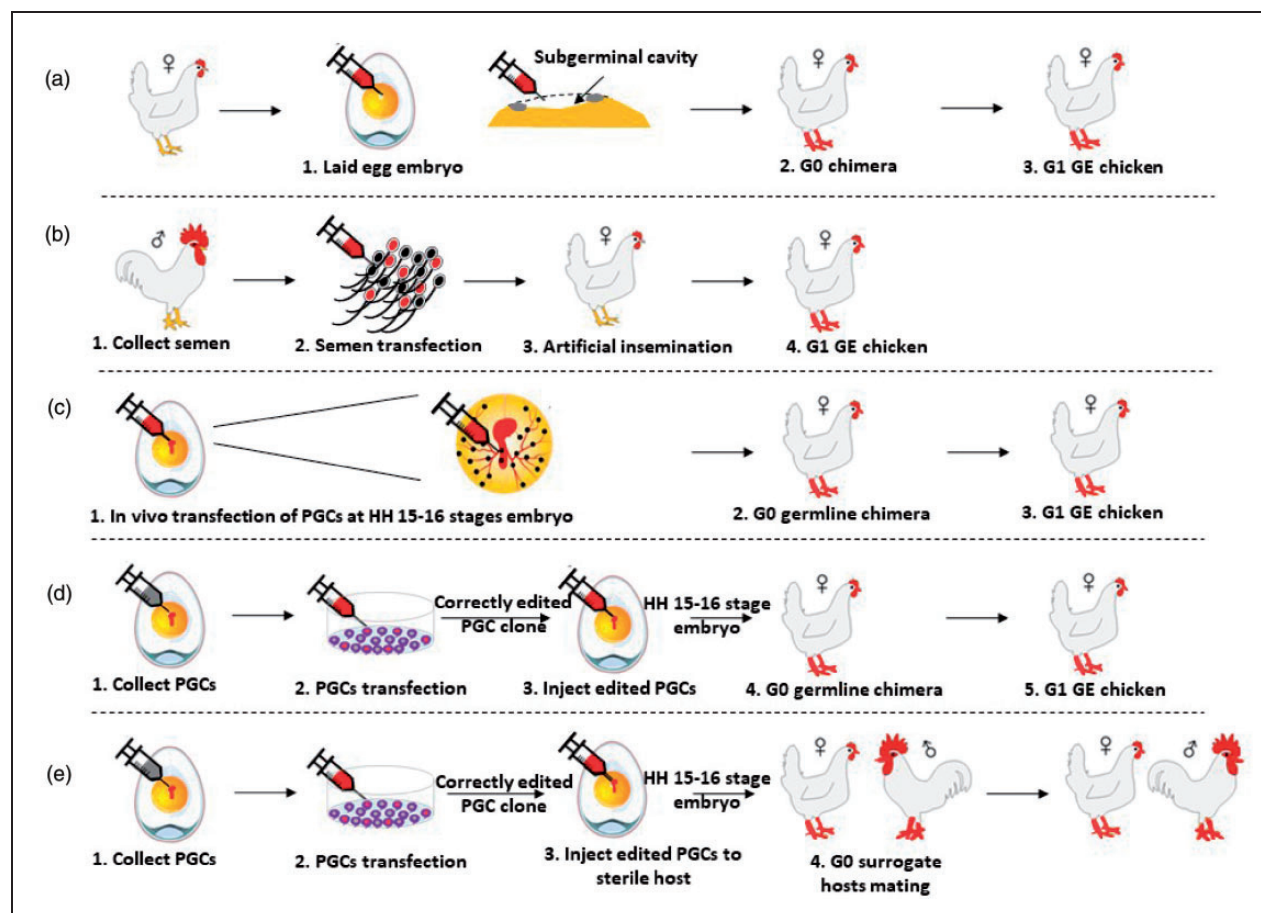


Figure 2. Schematic representation of different possible methods of establishing genome edited (GE) chicken. [a–e] the workflow of GE in the chicken via transduction with GE reagents to the embryonic blastoderm of a laid egg chicken embryo (a), sperm transfection for artificial insemination (b), primordial germ cell (PGC) transfection *in ovo* of embryos (c), PGC transfection *in vitro* culture (d), and transfer of *in vitro* GE PGCs to sterile host and subsequent mating of sterile G0 founders to generate pure GE chicken (e). After Looi et al.⁶⁴

embryo.^{28–30} In an improvement of this technique, Bartsch et al. combined Cas9 transgenic chicken embryos with the *in ovo* electroporation of guide RNAs, demonstrating *in ovo* electroporation in the chicken embryo is a feasible method of generating *in ovo* GE chickens.³¹ Similar experiments carried out in mammalian species require surgery for the *in utero* manipulations of the embryo with concomitant increased welfare impacts on the mother. Thus, the use of avian embryos represents a refinement by reducing the number of animals used and carrying out experiments at unprotected stages.

Blastoderm injection

Several research groups contributed the first genetically modified chickens using blastoderm injection of viral vectors.³² A small cavity lies under the blastoderm, the ‘subgerminal cavity’, into which viral vectors can be introduced which will then transduce the overlying

blastodermal cells and blastodermal germ cells (Figure 2(a)). The first genetically modified chicken was generated by the integration of the retroviral vector, known as avian leucosis virus (ALV), into the germline after blastoderm injection.³³ Later, the efficient generation of genetically modified chickens was reported using various replication-deficient viral vectors.^{34–36} Our laboratory and others showed an improved germline transmission of transgenes in chicken, quail and zebra finch using lentiviral-based vectors.^{36–39} Recently, adenoviruses were used to deliver a Cas9 transgene and a guide RNA directly to the quail blastoderm.^{40,41} Initially, authors targeted the quail melanophilin gene, resulting in grey plumage of homozygous GE quail offspring produced from the founder birds, whereas heterozygous and wild type quail exhibited dark brown plumage.⁴⁰ In 2020, the same laboratory targeted the quail myostatin gene, generated germline chimera and offspring. Homozygous myostatin edited quail showed significantly enhanced body

weight and muscle mass.⁴¹ In spite of the success of virus-mediated GE in quails and the feasibility of applying this delivery method of genome editors to other avian species, the efficiency of this virus-mediated method is low because of the low efficacy of transducing the blastodermal primordial germ cells after viral injection into the subgerminal cavity. In addition, increasing the size of lentiviral vectors to introduce Cas9 and the RNA guide leads to inefficient viral packaging that results in a low viral titer and reduced transduction efficacy. Similarly, windowing avian eggs for injection of virus followed by the subsequent hatching of chicks, will have varying survivability in different bird species.

Embryonic vascular system injection

A method for producing both genetically modified chicken and quail is by the direct injection of Tol2 transposon vectors into the vascular system of young embryos to target the germ cells during their migration through the circulation (Figure 2(c)).^{42,43} Transposons are self-integrating DNA vectors that can carry a transgenic cargo within the vector. Serralbo et al. succeeded in producing many fluorescent reporter lines of transgenic quail using this technique.⁴⁴ A limitation of this technique is that the transmission rate (frequency of producing GE offspring) is $\sim 1\%$. Recently, Challagulla et al. reported an *in ovo* delivery of GE components such as TALENs and CRISPR/Cas9, known as ‘*in vivo* transfection of PGCs [primordial germ cells]’, into the bloodstream of early embryos to generate GE offspring.⁴⁵ A limitation of this technique is the low germline transmission rate, $\sim 0.3\%$. Though this method indicates that the direct genome editing of any avian species is theoretically possible, the low transmission rate due to the low efficiency of genetically modifying the migratory PGCs in the vascular system and the resulting stable integration of the Cas9 vectors into the genome of the injected animal are drawbacks of this technique.

In vitro propagated avian PGCs

As stated above, germline-competent ES cells have not been isolated in avian species, which can be attributed to the early segregation of the germ cell lineage from the somatic cell lineage in birds. As is true for all vertebrate species, PGCs of birds are the progenitors or precursor cells of the sperm and oocytes. In freshly laid chicken eggs, as mentioned above, there are approximately 50 PGCs located within the centre of the blastoderm which contains approximately 40,000–60,000 cells.⁴⁶ Subsequently, PGCs migrate to the anterior germinal crescent from where 100–200 PGCs enter the

embryonic circulatory system at 48–60 h of incubation before migration to the forming gonad. PGCs will undergo sexual differentiation into spermatogonial stem cells in the male, producing spermatozoa. In females, PGCs undergo meiosis and will form mature oocytes upon sexual maturity.⁴⁷ In striking contrast to mammals, PGCs from chicken embryos can be isolated and cultured indefinitely, while keeping their commitment to the germline and germline competency.⁴⁸ Chicken PGCs are extracted from the vascular system at early developmental stages and placed in culture.⁴⁹ A defined serum-free culture medium for propagating chicken PGCs has been developed by optimising the signalling pathways necessary for avian germ cell self-renewal.⁵⁰

The long-term culture of PGCs does not compromise their ability to colonise the gonad when injected into the vascular system of surrogate host embryos and form functional gametes and offspring when the surrogate hosts are subsequently bred. This has been a turning point for the generation of GE chicken models through GE of culture PGCs (Figure 2(d) and (e)).⁵¹ Numerous reports based on using PGCs for the generation of GE chickens have been published including homologous recombination,⁵² TALENs,^{53,54} and CRISPR/Cas9 system.^{55,56} Among the genome editors, CRISPR/Cas9 has been successfully used in PGCs in developing functional genetic models by generating immunoglobulin heavy chain transgenic chicken,⁵⁵ ovomucoid KO chicken⁵⁶ and myostatin KO chicken.⁵⁷

Using cultured PGCs, a demonstration of GE for avian disease resistance has been achieved. Acidic nuclear phosphoprotein 32 family member A (ANP32A) was recently identified as a cellular host protein in birds required for avian influenza virus (IAV) polymerase adaptation and activity.⁵⁸ Long et al. used CRISPR/Cas9 to knock out the ANP32A in chicken PGCs, demonstrated that fibroblasts derived from GE PGCs did not support either mammalian or avian influenza virus polymerase activity during *in vitro* challenge experiments and were resilient to IAV infection.⁵⁹ Similarly, Koslová et al. and Hellmich et al. used CRISPR/Cas9 to delete the tryptophan residue number 38 of the Na⁺/H⁺ exchanger type 1 gene in PGCs, which is a critical amino acid for the entry of ALV subgroup J (ALV-J). The subsequent chickens generated from the GE PGCs were ALV-J resistant in *in vitro* and *in vivo* challenge experiments.^{60,61} The frequency of generating GE offspring from embryonic injections of GE PGCs is variable and usually from 1–50%.

Sperm mediated GE

Cooper et al. demonstrated that spermatozoa can be directly transfected and targeted with CRISPR/Cas9

GE tools.⁶² Subsequently, the GE sperm can be used for artificial insemination. This process is called sperm assisted gene editing (STAGE) and has been used to make GE chickens in a single generation (Figure 2(b)). However, the drawbacks of this technique are the genetic deletions in the offspring did not closely correspond to the cleavage site of the CRISPR/Cas9 in the genome and the rate of genetic modification in the offspring from the founders was from 0–26%.⁶²

The above results show that genetic modification of chicken and quail is highly advanced. A comprehensive list of genetically modified chicken and quail available in different research laboratories are reported in other reviews.^{44,63}

Use of avian sterile hosts to address the principles of the 3Rs

A key challenge for GE of bird species is to efficiently produce GE offspring from the mosaic founder or surrogate hosts without breeding hundreds of non-GE offspring. Until now, the most successful method of generating genome edited chickens is by injecting exogenous edited PGCs intravenously into stage 15–16⁺ HH (Hamburger Hamilton) surrogate embryo in windowed eggs. However, the edited donor PGCs and endogenous PGCs compete with each other to form functional gametes. Thus, there is a greater opportunity for the endogenous PGCs to transmit its genome to the subsequent generation of offspring than the donor PGCs. To expedite the germline transmission from the donor edited PGCs, endogenous PGCs can be eliminated by busulfan and γ -irradiation, both of which can cause death and health problems for the treated host bird.^{65,66} However, GE can be used to create gene knockouts to eliminate the competition from endogenous PGCs by rendering the host bird devoid of sperm or eggs. Two sterile chicken host models have been produced: a TALEN-mediated knockout of the germ cell determinant, *DDX4* (DEAD-Box helicase 4) which resulted in sterile hens, and CRISPR/Cas9 insertion of an inducible transgene insertion (iCaspase9) which generated sterile cockerels and hens. Both models transmitted only donor PGCs to the next generation offspring.^{53,54,67} Remarkably, the direct mating of G0 sterile host cockerels and hens carrying GE donor PGCs leads to the generation of pure GE offspring in one generation. This technique is described as ‘sire dam surrogate mating’ (Figure 2(e)). Thus, sterile chicken surrogate hosts aid in the generation of GE birds using a reduced number of animals, thus supporting 3Rs principles, and is potentially applicable in projects for the ‘cryoconservation’ of poultry and endangered avian breeds.

The future for GE in other avian species

The injection of GE reagents directly into the bloodstream of embryonated eggs through a window made in the egg shell can theoretically be applied to all bird species. However, this technique is inefficient for the generation of transgenic and GE offspring (0.3–1%) and the hatch rate of the windowed injected eggs will differ drastically between bird species.^{68,69} Transmission rates of the genetic modification using this technique are currently low in chicken and multiple generations will be needed to generate pure offspring containing a homozygous GE allele. These methods will prove more difficult when applied to other avian model species such as zebra finch, ducks and turkeys. It would be extremely beneficial to be able to propagate the PGCs *in vitro* for other avian species and to develop sterile surrogate birds that would be appropriate hosts for the germ cells from multiple species.^{70–72} It has been demonstrated that male germ cells can form functional gametes when transplanted between evolutionarily distant bird species such as chicken to duck, pheasant to chicken, chicken to guinea fowl, and chicken to the houbara bustard.^{68,73–75} Thus, it should be possible to use a sterile chicken host for male genome modified PGCs from any bird species. However, oocyte and egg development has not been demonstrated after PGC transplantation between bird species. The oocyte size varies greatly between bird species and the somatic granulosa cells that surround the oocyte control both oocyte maturation and ovulation. It remains to be determined if hybrid follicles comprised of donor germ cells from one bird species can form a functional egg in another bird species.

Conclusions

The chicken has been a mainstay of vertebrate embryology research for many decades because their embryos come conveniently packaged in eggs and offer easy accessibility of developmental stages for experimental manipulations.¹¹ Many fundamental questions yet have to answer in developmental biology around neural development, organogenesis and patterning of the embryo. The advancement of the GE tools in chicken in combination with live-cell imaging and single-cell transcriptomics in both embryos and adult chicken has the potential of exploring these pertinent questions. Transferring these tools to other bird species would facilitate research efforts in many laboratory settings.

The chicken is an invaluable model for studying basic immunology and provides seminal contributions to fundamental immunological principles such as graft-versus-host responses and the key role of lymphocytes in adaptive immunity.⁷⁶ Infectious disease outbreaks in

poultry is a persistent threat to the poultry sector as well as concerns of zoonotic transmission. GE tools can help to investigate disease-resistance in birds and also heat tolerance to combat climate impact on poultry production. In summary, the ongoing improvements in genome modification tools with avian embryo manipulation and PGC culture continues to refine the production of GE avian species with positive outputs on the 3R principles.

Author contributions

Conceptualisation, SKP and MJM; writing – original draft preparation, SKP; writing – review and editing, SKP and MJM; visualisation, SKP; supervision, MJM; funding acquisition, MJM.


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Résumé

Les espèces aviaires sont utilisées comme systèmes modèles dans la recherche et ont contribué à des concepts révolutionnaires en matière de biologie du développement, d'immunologie, de génétique, de virologie, de cancer et de biologie cellulaire. Le poulet en particulier est un modèle de recherche important ainsi qu'un animal agricole constituant une source majeure de protéines animales pour la population mondiale. Le développement de méthodes d'édition du génome, y compris CRISPR/Cas9, permettant de médier l'ingénierie de lignée germinale du génome aviaire, aura des applications importantes dans les activités biomédicales, agricoles et biotechnologiques. Ces outils précis d'édition du génome ont notamment le potentiel d'améliorer la santé et la productivité des oiseaux en identifiant et en validant les variantes génétiques bénéfiques chez les populations d'oiseaux. Nous présentons ici une description concise des méthodes existantes et des applications actuelles des outils d'édition du génome chez les espèces d'oiseaux, axés sur les poulets, en mettant l'accent sur l'utilisation des animaux et les questions de bien-être pour chacune des techniques présentées.

Abstract

Vogelarten dienen in der Forschung als Modellsysteme und haben zu bahnbrechenden Konzepten in der Entwicklungsbiologie, Immunologie, Genetik, Virologie, Krebsforschung und Zellbiologie beigetragen. Insbesondere das Huhn, das als Nutztier einen großen Beitrag zur Versorgung der Weltbevölkerung mit tierischem Protein leistet, ist ein wichtiges Forschungsmodell. Die Entwicklung von Genome-Editing-Methoden, darunter CRISPR/Cas9, für Keimbahn-Engineering des aviären Genoms wird zu wichtigen Anwendungen in der Biomedizin, Landwirtschaft und Biotechnologie führen. Insbesondere haben diese präzisen Genome-Editing-Tools das Potenzial, die Gesundheit und Produktivität von Vögeln zu verbessern, indem vorteilhafte genetische Varianten in Vogelpopulationen identifiziert und validiert werden. Hier präsentieren wir eine kurze Beschreibung der existierenden Methoden und aktuellen Anwendungen der Genome-Editing-Tools bei Vogelarten, wobei der Schwerpunkt auf Hühnern liegt. Besonderes Augenmerk gilt Fragen der Tiernutzung und des Tierschutzes für die einzelnen vorgestellten Techniken.

Resumen

Las especies aviares se utilizan como sistemas de modelos de animales para la investigación y han contribuido en conceptos innovadores de la biología del desarrollo, en la inmunología, la genética, la virología, la lucha contra el cáncer y la biología celular. El pollo es especialmente un modelo de investigación y animal agrícola de gran importancia ya que representa un gran contribuidor para las fuentes de proteínas de animales para la población global. El desarrollo de los métodos de edición genómica, incluido CRISPR/Cas9, para mediar la ingeniería la línea germinal del genoma aviar tendrá aplicaciones importantes en las distintas actividades biomédicas, agrícolas y biotecnológicas. En especial, estas herramientas de edición genómica precisas tienen el potencial de mejorar la productividad y la salud aviar identificando y validando variantes genéticas beneficiosas en las poblaciones aviares. En este estudio presentamos una descripción concisa de los métodos existentes y las aplicaciones actuales de las herramientas de edición genómica en especies aviares, centrándonos en pollos y prestando atención al uso y el bienestar animal para cada una de las técnicas presentadas.